

#8

J620 WA



PCT10

RAW SEQUENCE LISTING

DATE: 11/18/2002

PATENT APPLICATION: US/10/089,787

TIME: 15:15:14

Input Set : A:\20522p.txt

Output Set: N:\CRF4\11182002\J089787.raw

4 <110> APPLICANT: El-Sherbeini, Mohamed
 5 Azzolina, Barbara A.
 7 <120> TITLE OF INVENTION: MraY GENE AND ENZYME OF PSEUDOMONAS
 8 AERUGINOSA
 10 <130> FILE REFERENCE: 20522P
 12 <140> CURRENT APPLICATION NUMBER: 10/089,787
 13 <141> CURRENT FILING DATE: 2002-04-03
 15 <150> PRIOR APPLICATION NUMBER: PCT/US00/27056
 16 <151> PRIOR FILING DATE: 2000-09-29
 18 <150> PRIOR APPLICATION NUMBER: 60/157,580
 19 <151> PRIOR FILING DATE: 1999-10-04

21 <160> NUMBER OF SEQ ID NOS: 4
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0

25 <210> SEQ ID NO: 1

26 <211> LENGTH: 1083

27 <212> TYPE: DNA

28 <213> ORGANISM: Pseudomonas Aeruginosa

30 <400> SEQUENCE: 1

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32 tacctgaccc tgcgggcat tctcagcgtg ctcaccgcgc tgtcgctgtc gctgtggctg	120
33 gggccctgga tgatccgtac cttgcagatc ccccgatcg gccaggccgt gcgcaacgac	180
34 ggtccgcagt cgcacctgtc gaagaagggc accccgacca tgggcggcgc cctgatcctt	240
35 accgccatag ccatcagcac gctgctgtgg gcggatcttt ccaaccgcta cgtgtgggta	300
36 gtgctggctg ttacctgtct gttcggtgcc atcggctggg tagacgacta ccgcaagggtg	360
37 atcgagaaga actcccgtag cctgccgagc cgctggaagt acttctggca gtcggtgttc	420
38 ggcacggcgc ccgccgtgtt cctctacatg actgccgaaa ccccgatcga gaccaccctg	480
39 atcgtgccga tgctgaagag cgtcgagatc cagttgggca tcttcttcgt ggtcctgacc	540
40 tacttcgtca tcgtcggtct gagcaatgcg gtgaacctca ccgacggtct cgacggcctg	600
41 gcgatcatgc cgacggtaat ggttgccggc gcgctgggca tcttctgcta cctgtcgggc	660
42 aacgtgaagt tcgccgagta cctgctgatt cccaacgtac cgggcgccgc cgagctgatc	720
43 gtgttctgcg ccgcgctggt cgccgccggc ctccgcttcc tctggttcaa cacctatccg	780
44 gcgcaggctc tcatgggcga cgtcggcgcg ctggcgctgg gcgccgcgct gggcaccatc	840
45 gcggtgatcg tgcgccagga gatcgtgctg ttcacatggt gtggggtgtt cgtcatggaa	900
46 accctctcgg tgatgatcca ggtcgcttcc ttcaagctga ccggacgccg cgtcttccgt	960
47 atggcgccga tccatcacca tttcgaactg aaaggctggc cggacccgcg cgtgatcgtg	1020
48 cgcttctgga tcatcaccgt gatcctggtg ctgatcggcc tcgccacctt gaagctgcgt	1080
49 tga	1083

51 <210> SEQ ID NO: 2

52 <211> LENGTH: 360

53 <212> TYPE: PRT

54 <213> ORGANISM: Pseudomonas Aeruginosa

56 <400> SEQUENCE: 2

57 Met Leu Leu Leu Leu Ala Glu Tyr Leu Gln Gln Phe Tyr Lys Gly Phe

ENTERED

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58 1          5          10          15
59 Gly Val Phe Gln Tyr Leu Thr Leu Arg Gly Ile Leu Ser Val Leu Thr
60          20          25          30
61 Ala Leu Ser Leu Ser Leu Trp Leu Gly Pro Trp Met Ile Arg Thr Leu
62          35          40          45
63 Gln Ile Pro Gln Ile Gly Gln Ala Val Arg Asn Asp Gly Pro Gln Ser
64          50          55          60
65 His Leu Ser Lys Lys Gly Thr Pro Thr Met Gly Gly Ala Leu Ile Leu
66 65          70          75          80
67 Thr Ala Ile Ala Ile Ser Thr Leu Leu Trp Ala Asp Leu Ser Asn Arg
68          85          90          95
69 Tyr Val Trp Val Val Leu Val Val Thr Leu Leu Phe Gly Ala Ile Gly
70          100          105          110
71 Trp Val Asp Asp Tyr Arg Lys Val Ile Glu Lys Asn Ser Arg Gly Leu
72          115          120          125
73 Pro Ser Arg Trp Lys Tyr Phe Trp Gln Ser Val Phe Gly Ile Gly Ala
74          130          135          140
75 Ala Val Phe Leu Tyr Met Thr Ala Glu Thr Pro Ile Glu Thr Thr Leu
76 145          150          155          160
77 Ile Val Pro Met Leu Lys Ser Val Glu Ile Gln Leu Gly Ile Phe Phe
78          165          170          175
79 Val Val Leu Thr Tyr Phe Val Ile Val Gly Ser Ser Asn Ala Val Asn
80          180          185          190
81 Leu Thr Asp Gly Leu Asp Gly Leu Ala Ile Met Pro Thr Val Met Val
82          195          200          205
83 Ala Gly Ala Leu Gly Ile Phe Cys Tyr Leu Ser Gly Asn Val Lys Phe
84          210          215          220
85 Ala Glu Tyr Leu Leu Ile Pro Asn Val Pro Gly Ala Gly Glu Leu Ile
86 225          230          235          240
87 Val Phe Cys Ala Ala Leu Val Gly Ala Gly Leu Gly Phe Leu Trp Phe
88          245          250          255
89 Asn Thr Tyr Pro Ala Gln Val Phe Met Gly Asp Val Gly Ala Leu Ala
90          260          265          270
91 Leu Gly Ala Ala Leu Gly Thr Ile Ala Val Ile Val Arg Gln Glu Ile
92          275          280          285
93 Val Leu Phe Ile Met Gly Gly Val Phe Val Met Glu Thr Leu Ser Val
94          290          295          300
95 Met Ile Gln Val Ala Ser Phe Lys Leu Thr Gly Arg Arg Val Phe Arg
96 305          310          315          320
97 Met Ala Pro Ile His His His Phe Glu Leu Lys Gly Trp Pro Asp Pro
98          325          330          335
99 Arg Val Ile Val Arg Phe Trp Ile Ile Thr Val Ile Leu Val Leu Ile
100          340          345          350
101 Gly Leu Ala Thr Leu Lys Leu Arg
102          355          360
104 <210> SEQ ID NO: 3
105 <211> LENGTH: 29
106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence

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109 <220> FEATURE:
110 <223> OTHER INFORMATION: PCR Primer
112 <400> SEQUENCE: 3
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116 <211> LENGTH: 27
117 <212> TYPE: DNA
118 <213> ORGANISM: Artificial Sequence
120 <220> FEATURE:
121 <223> OTHER INFORMATION: PCR Primer
123 <400> SEQUENCE: 4
124 ttggatcctc aacgcagctt caagggtg 27

VERIFICATION SUMMARY

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